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mask repetitive contig sequence unique

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Analysis of major **repetitive** DNA sequences in the dog(*Canis familiaris*) genome

S Bentolila, JM Bach, JL Kessler, I Bordelais & ... - Mammalian Genome, 1999 - springerlink.com

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CV Jongeneel - Briefings in Bioinformatics, 2000 - ingentaconnect.com

... from several species, under the denomination of **unique** gene indices. ... to return gapped alignments between the query and the database **sequence**. ... **Contig** databases ...Cited by 12 - [Web Search](#) - [ismb00.sdsc.edu](#) - [sib-dea.unil.ch](#) - [amber.cs.umd.edu](#) - [all 9 versions »](#)Gene discovery by EST sequencing in *Toxoplasma gondii* reveals sequences restricted to the

JW Ajioka, JC Boothroyd, BP Brunk, A Hehl, L ... - Genome Res, 1998 - genome.org

... As in (3), above, **repetitive** elements were identified ... Lipman, unpubl.) was used to**mask** low entropy ... **sequence** was determined for each **contig** defined (available ...Cited by 75 - [Web Search](#) - [genome.org](#) - [path.cam.ac.uk](#) - [ncbi.nlm.nih.gov](#) - [all 7 versions »](#)Comparative Genome Mapping in the **Sequence**-based Era: Early Experience with Human Chromosome 7

JW Thomas, TJ Summers, SQ Lee-Lin, VV Maduro, JR ... - Genome Research, 2000 - genome.org

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H Nagano, L Wu, S Kawasaki, Y Kishima, Y Sano - Genome, 1999 - article.pubs.nrc-cnrc.gc.ca

... repeti- tive short **sequence**(s), which can **mask** their own ... the single-copy regions in the 260-kb **contig** are interrupted by **repetitive** sequences every ...Cited by 7 - [Web Search](#) - [dx.doi.org](#) - [pubs.nrc-cnrc.gc.ca](#) - [ncbi.nlm.nih.gov](#) - [all 6 versions »](#)The Mouse Clock Locus: **Sequence** and Comparative Analysis of 204 Kb from Mouse Chromosome 5

LD Wilsbacher, AM Sangoram, MP Antoch, JS ... - Genome Research, 2000 - genome.org

... 11-22 align within the largest **contig**, while exons ... of introns as well (mouse **repetitive** elements were ... is possible that position effects could **mask** a regulatory ...Cited by 4 - [Web Search](#) - [pubmedcentral.nih.gov](#) - [dx.doi.org](#) - [ncbi.nlm.nih.gov](#) - [all 6 versions »](#)Identification of a gene that reverses the immortal phenotype of a subset of cells and is a member ...

MJ Bertram, NG Berube, X Hang-Swanson, Q Ran, JK ... - Mol. Cell. Biol, 1999 - mcb.asm.org

... cDNAs containing **repetitive** elements were eliminated from further ... have coding **sequence** within the **contig** (1). One ... cell in the population to **mask** the senescence ...Cited by 60 - [Web Search](#) - [mcb.asm.org](#) - [pubmedcentral.nih.gov](#) - [ncbi.nlm.nih.gov](#) - [all 5 versions »](#)Data mining parasite genomes: haystack searching with a computer

D LAWSON - Parasitology, 1999 - journals.cambridge.org

... The generation of assembled **contig** consensus from the primary ... regions of low- complexity can **mask** true hits ... genome or the degree of **repetitive** sequences present ...
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K Hamada, T Kohno, M Takahashi, M Yamazaki, H ... - Genes Chromosomes and Cancer, 2000 - [doi.wiley.com](#)
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The search for Lps: 1993-1998

B Beutler, A Poltorak - Journal of Endotoxin Research, 2000 - [ingentaconnect.com](#)
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D Rollinson - 1999 - [print.google.com](#)
 ... A genome **sequence** lists the **unique** order of all nucleotides and their arrangement into coding and non-coding regions and is essentially a starting point for ...
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RE Matick - IBM Systems Journal, 1986 - [research.ibm.com](#)
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[View as HTML](#) - [Web Search](#) - [domino.research.ibm.com](#) - [csa.com](#)

Free Software in Molecular Biology for Macintosh and MS Windows Computers

D Gilbert - Bioinformatics Methods and Protocols (S. Misener and SA ..., 1999 - [iubio.bio.indiana.edu](#)
 ... a MS DOS program to identify common **repetitive** elements in ... also able to identify and **mask** vector **sequence** ... ClustalW multiple alignment, CAP **contig** assembly, tacg ...
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D Huertas, S Howe, A McGuigan, C Huxley - Human Molecular Genetics, 2000 - [hmg.oupjournals.org](#)
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SR Wolman, S Sell - 1997 - [print.google.com](#)

... For many forms of human cancer, however, **unique** and tumor-specific chromosomal patterns have not yet been discerned, largely because of the extent and ...

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L3	9	mask and repetitive and contig and sequence and unique and @ad<"20000101"	USPAT	OR	OFF	2005/12/01 14:32
L4	7	mask and repetitive and contig and sequence and unique and @ad<"20000101" and database	USPAT	OR	OFF	2005/12/01 14:32